



## SEQUENCE LISTING

<10> Endress, Gregory A.  
Rosen, Craig A.

<120> Prostate Specific Secreted Protein

<130> PF457

<140> Unassigned

<141> 1999-03-30

<150> 60/080,311

<151> 1998-04-01

<150> 60/080,898

<151> 1998-04-07

<160> 15

<170> PatentIn Ver. 2.0

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<213> Homo sapiens

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cccagtcaca ggcgagagcc ctggg atg cac cgg cca gag gcc atg ctg ctg 172
                               Met His Arg Pro Glu Ala Met Leu Leu
                               1           5
ctg ctc acg ctt gcc ctc ctg ggg ggc ccc acc tgg gca ggg aag atg 220
Leu Leu Thr Leu Ala Leu Leu Gly Gly Pro Thr Trp Ala Gly Lys Met
10           15           20           25
tat ggc cct gga gga ggc aag tat ttc agc acc act gaa gac tac gac 268
Tyr Gly Pro Gly Gly Gly Lys Tyr Phe Ser Thr Thr Glu Asp Tyr Asp
           30           35           40
cat gaa atc aca ggg ctg cgg gtg tct gta ggt ctt ctg ctg gtg aaa 316
His Glu Ile Thr Gly Leu Arg Val Ser Val Gly Leu Leu Leu Val Lys
           45           50           55
agt gtc cag gtg aaa ctt gga gac tcc tgg gac gtg aaa ctg gga gcc 364
Ser Val Gln Val Lys Leu Gly Asp Ser Trp Asp Val Lys Leu Gly Ala
           60           65           70
tta ggt ggg aat acc cag gaa gtc acc ctg cag cca ggc gaa tac atc 412
Leu Gly Gly Asn Thr Gln Glu Val Thr Leu Gln Pro Gly Glu Tyr Ile
           75           80           85
aca aaa gtc ttt gtc gcc ttc caa gct ttc ctc cgg ggt gtg gtc atg 460
Thr Lys Val Phe Val Ala Phe Gln Ala Phe Leu Arg Gly Val Val Met
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tac acc agc aag gac cgc tat ttc tat ttt ggg aag ctt gat ggc cag 508
Tyr Thr Ser Lys Asp Arg Tyr Phe Tyr Phe Gly Lys Leu Asp Gly Gln
           110           115           120
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Ile Ser Ser Ala Tyr Pro Ser Gln Glu Gly Gln Val Leu Val Gly Ile
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125                      130                      135  
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 Tyr Gly Gln Tyr Gln Leu Leu Gly Ile Lys Ser Ile Gly Phe Glu Trp  
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 aat tat cca cta gag gag ccg acc act gag cca cca gtt aat ctc aca 652  
 Asn Tyr Pro Leu Glu Glu Pro Thr Thr Glu Pro Pro Val Asn Leu Thr  
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 tac tca gca aac tca ccc gtg ggt cgc tag ggtggggtat ggggccatcc 702  
 Tyr Ser Ala Asn Ser Pro Val Gly Arg  
       170                      175  
 gagctgaggc catctgggtg gtggtggctg atggtactgg agtaactgag tcgggacgct 762  
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                     20                      25                      30  
 Tyr Phe Ser Thr Thr Glu Asp Tyr Asp His Glu Ile Thr Gly Leu Arg  
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 Val Ser Val Gly Leu Leu Leu Val Lys Ser Val Gln Val Lys Leu Gly  
                     50                      55                      60  
 Asp Ser Trp Asp Val Lys Leu Gly Ala Leu Gly Gly Asn Thr Gln Glu  
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 Val Thr Leu Gln Pro Gly Glu Tyr Ile Thr Lys Val Phe Val Ala Phe  
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 Phe Tyr Phe Gly Lys Leu Asp Gly Gln Ile Ser Ser Ala Tyr Pro Ser  
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 Gln Glu Gly Gln Val Leu Val Gly Ile Tyr Gly Gln Tyr Gln Leu Leu  
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 Gly Ile Lys Ser Ile Gly Phe Glu Trp Asn Tyr Pro Leu Glu Glu Pro  
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Asn Asn Ile Leu Ser Ser Ile Arg Leu Lys Phe Gly Asn Asn Trp Ser			
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Gln Glu Tyr Gly Ser Ser Gly Arg Ala Glu Ile Glu Val Lys Leu Asn			
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Pro Asp Glu Thr Val Leu Gly Phe Ser Gly Ser Phe Tyr Ile Phe Met			
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His Gln Ile Ile Ile Thr Thr Ser Gln Pro Arg Glu Leu Ile Ile Gly			
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Pro Leu Thr Gly Arg Tyr Val Tyr Thr Ser Tyr Pro Glu Asn Pro Asn			
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 tctcccgga tcttgaggtc acatgcgtgg tgggtggacgt aagccacgaa gaccctgagg 180  
 tcaagttcaa ctggtacgtg gacggcgtgg aggtgcataa tgccaagaca aagccgcggg 240  
 aggagcagta caacagcacg taccgtgtgg tcagcgtcct caccgtcctg caccaggact 300  
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 catcccggga tgagctgacc aagaaccagg tcagcctgac ctgcctggtc aaaggcttct 480  
 atccaagcga catgccgtg gagtgggaga gcaatgggca gccggagaac aactacaaga 540  
 ccacgcctcc cgtgctggac tccgacggt ccttcttct ctacagcaag ctcaccgtgg 600  
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 gccctaact ccgcccagtt ccgcccattc tccgccccat ggctgactaa ttttttttat 180  
 ttatgcagag gccgaggccg cctcggcctc tgagctatlc cagaagtagt gaggaggctt 240  
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 cagttccgcc cattctccgc cccatggctg actaattttt tttatttatg cagaggccga 180  
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